



## REPLACEMENT SHEET 10/800,449

Figure 2A. DNA and deduced amino acid sequences of hG-CSF-L-vFc<sub>12</sub>

|   |               |
|---|---------------|
| DNA   | SEQ ID NO: 17 |
| Amino Acid Sequence   | SEQ ID NO: 18 |
| aag ctt ccc aga ccc atg gct gga cct gcc acc cag agc ccc atg aag ctg atg gcc ctg<br>HindIII M A G P A T Q S P M K L M A L 60     |               |
| -30 -20   |               |
| cag ctg ctg ctg tgg cac agt gca ctc tgg aca gtg cag gaa gcc acc ccc ctg ggc cct 120<br>Q L L L W H S A L W T V Q E A T P L G P  |               |
| -10 -1 1  |               |
| gcc agc tcc ctg ccc cag agc ttc ctg ctc aag tgc tta gag caa gtg agg aag atc cag 180<br>A S S L P Q S F L L K C L E Q V R K I Q  |               |
| 10 20   |               |
| ggc gat ggc gca gcg ctc cag gag aag ctg tgt gcc acc tac aag ctg tgc cac ccc gag 240<br>G D G A A L Q E K L C A T Y K L C H P E  |               |
| 30 40   |               |
| gag ctg gtg ctg ctc gga cac tct ctg ggc atc ccc tgg gct ccc ctg agc agc tgc ccc 300<br>E L V L L G H S L G I P W A P L S S C P  |               |
| 50 60   |               |
| agc cag gcc ctg cag ctg gca ggc tgc ttg agc caa ctc cat agc ggc ctt ttc ctc tac 360<br>S Q A L Q L A G C L S Q L H S G L F L Y  |               |
| 70 80   |               |
| cag ggg ctc ctg cag gcc ctg gaa ggg atc tcc ccc gag ttg ggt ccc acc ttg gac aca 420<br>Q G L L Q A L E G I S P E L G P T L D T  |               |
| 90 100  |               |
| ctg cag ctg gac gtc gcc gac ttt gcc acc acc atc tgg cag cag atg gaa gaa ctg gga 480<br>L Q L D V A D F A T T I W Q Q M E E L G  |               |
| 110 120   |               |
| atg gcc cct gcc ctg cag ccc acc cag ggt gcc atg ccg gcc ttc gcc tct gct ttc cag 540<br>M A P A L Q P T Q G A M P A F A S A F Q  |               |
| 130 140   |               |
| cgc cgg gca gga ggg gtc cta gtt gcc tcc cat ctg cag agc ttc ctg gag gtg tgc tac 600<br>R R A G G V L V A S H L Q S F L E V S Y  |               |
| 150 160   |               |
| cgc gtt cta cgc cac ctt gcc cag ccc gga tcc ggt ggc ggt tcc ggt gga ggc gga agc 660<br>R V L R H L A Q P G S G G G S G G G S    |               |
| 170 180   |               |
| ggc ggt gga gga tca gag cgc aaa tgt tgt gtc gag tgc cca ccg tgc cca gca cca cct 720<br>G G G G S E R K C C V E C P P C P A P P  |               |
| 190 200   |               |
| gtg gca gga ccg tca gtc ttc ctc ttc ccc cca aaa ccc aag gac acc ctc atg atc tcc 780<br>V A G P S V F L F P P K P K D T L M I S  |               |
| 210 220   |               |
| cgg acc cct gag gtc acg tgc gtg gtg gtg gac gtg agc cac gaa gac ccc gag gtc cag 840<br>R T P E V T C V V V D V S H E D P E V Q  |               |
| 230 240   |               |
| ttc aac tgg tac gtg gac ggc gtg gag gtg cat aat gcc aag aca aag cca cgg gag gag 900<br>F N W Y V D G V E V H N A K T K P R E E  |               |
| 250 260   |               |
| cag ttc aac agc acg ttc cgt gtg gtc agc gtc ctc acc gtt gtg cac cag gac tgg ctg 960<br>Q F N S T F R V V S V L T V V H Q D W L  |               |
| 270 280   |               |
| aac ggc aag gag tac aag tgc aag gtc tcc aac aaa ggc ctc cca gcc tcc atc gag aaa 1020<br>N G K E Y K C K V S N K G L P A S I E K |               |
| 290 300   |               |
| acc atc tcc aaa acc aaa ggg cag ccc cga gaa cca cag gtg tac acc ctg ccc cca tcc 1080<br>T I S K T K G Q P R E P Q V Y T L P P S |               |
| 310 320   |               |
| cgg gag gag atg acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa ggc ttc tac ccc 1140<br>R E E M T K N Q V S L T C L V K G F Y P |               |
| 330 340   |               |
| agc gac atc gcc gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc aca 1200<br>S D I A V E W E S N G Q P E N N Y K T T |               |
| 350 360   |               |
| cct ccc atg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc acc gtg gac aag 1260<br>P P M L D S D G S F F L Y S K L T V D K |               |
| 370 380   |               |
| agc agg tgg cag cag ggg aac gtc ttc tca tgc tcc gtg atg cat gag gct ctg cac aac 1320<br>S R W Q Q G N V F S C S V M H E A L H N |               |
| 390 400   |               |
| cac tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa tga gaa ttc 1368<br>H Y T Q K S L S L S P G K EcoRI                         |               |
| 410 418   |               |

**Figure 2B. DNA and deduced amino acid sequences of hG-CSF-L-vFc<sub>γ4</sub>**

|   |               |
|---|---------------|
| DNA   | SEQ ID NO: 19 |
| Amino Acid Sequence   | SEQ ID NO: 20 |
| aag ctt ccc aga ccc atg gct gga cct gcc acc cag agc ccc atg aag ctg atg gcc ctg<br>HindIII M A G P A T Q S P M K L M A L 60     |               |
| -30   | -20           |
| cag ctg ctg ctg tgg cac agt gca ctc tgg aca gtg cag gaa gcc acc ccc ctg ggc cct 120<br>Q L L L W H S A L W T V Q E A T P L G P  |               |
| -10   | -1            |
| gcc agc tcc ctg ccc cag agc ttc ctg ctc aag tgc tta gag caa gtg agg aag atc cag 180<br>A S S L P Q S F L L K C L E Q V R K I Q  |               |
| 10  | 20            |
| ggc gat ggc gca gcg ctc cag gag aag ctg tgt gcc acc tac aag ctg tgc cac ccc gag 240<br>G D G A A L Q E K L C A T Y K L C H P E  |               |
| 30  | 40            |
| gag ctg gtg ctg ctc gga cac tct ctg ggc atc ccc tgg gct ccc ctg agc agc tgc ccc 300<br>E L V L L G H S L G I P W A P L S S C P  |               |
| 50  | 60            |
| agc cag gcc ctg cag ctg gca ggc tgc ttg agc caa ctc cat agc ggc ctt ttc ctc tac 360<br>S Q A L Q L A G C L S Q L H S G L F L Y  |               |
| 70  | 80            |
| cag ggg ctc ctg cag gcc ctg gaa ggg atc tcc ccc gag ttg ggt ccc acc ttg gac aca 420<br>Q G L L Q A L E G I S P E L G P T L D T  |               |
| 90  | 100           |
| ctg cag ctg gac gtc gcc gac ttt gcc acc acc atc tgg cag cag atg gaa gaa ctg gga 480<br>L Q L D V A D F A T T I W Q Q M E E L G  |               |
| 110   | 120           |
| atg gcc cct gcc ctg cag ccc acc cag ggt gcc atg ccg gcc ttc gcc tct gct ttc cag 540<br>M A P A L Q P T Q G A M P A F A S A F Q  |               |
| 130   | 140           |
| cgc cgg gca gga ggg gtc cta gtt gcc tcc cat ctg cag agc ttc ctg gag gtg tgc tac 600<br>R R A G G V L V A S H L Q S F L E V S Y  |               |
| 150   | 160           |
| cgc gtt cta cgc cac ctt gcc cag ccc gga tcc ggt ggc ggt tcc ggt gga ggc gga agc 660<br>R V L R H L A Q P G S G G G S G G G G S  |               |
| 170   | 180           |
| ggc ggt gga gga gga gag tcc aaa tat ggt ccc cca tgc cca cca tgc cca gca cct gag 720<br>G G G G S E S K Y G P P C P P C P A P E  |               |
| 190   | 200           |
| ttc gag ggg gga cca tca gtc ttc ctg ttc ccc cca aaa ccc aag gac act ctc atg atc 780<br>F A G G P S V F L F P P K P K D T L M I  |               |
| 210   | 220           |
| tcc cgg acc cct gag gtc acg tgc gtg gtg gtg gac gtg agc cag gaa gac ccc gag gtc 840<br>S R T P E V T C V V V D V S Q E D P E V  |               |
| 230   | 240           |
| cag ttc aac tgg tac gtg gat ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg gag 900<br>Q F N W Y V D G V E V H N A K T K P R E  |               |
| 250   | 260           |
| gag cag ttc aac agc acg tac cgt gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg 960<br>E Q F N S T Y R V V S V L T V L H Q D W  |               |
| 270   | 280           |
| ctg aac ggc aag gag tac aag tgc aag gtc tcc aac aaa ggc ctc ccg tcc tcc atc gag 1020<br>L N G K E Y K C K V S N K G L P S S I E |               |
| 290   | 300           |
| aaa acc atc tcc aaa gcc aaa ggg cag ccc cga gag cca cag gtg tac acc ctg ccc cca 1080<br>K T I S K A K G Q P R E P Q V Y T L P P |               |
| 310   | 320           |
| tcc cag gag gag atg acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa ggc ttc tac 1140<br>S Q E E M T K N Q V S L T C L V K G F Y |               |
| 330   | 340           |
| ccc agc gac atc gcc gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc 1200<br>P S D I A V E W E S N G Q P E N N Y K T |               |
| 350   | 360           |
| acg cct ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc agg cta acc gtg gac 1260<br>T P P V L D S D G S F F L Y S R L T V D |               |
| 370   | 380           |
| aag agc agg tgg cag gag ggg aat gtc ttc tca tgc tcc gtg atg cat gag gct ctg cac 1320<br>K S R W Q E G N V F S C S V M H E A L H |               |
| 390   | 400           |
| aac cac tac aca cag aag agc ctc tcc ctg tct ctg ggt aaa tga gaa ttc 1371<br>N H Y T Q K S L S L S L G K EcoRI                   |               |
| 410   | 419           |

**Figure 2C. DNA and deduced amino acid sequences of hG-CSF-L-vFc<sub>γ1</sub>**

|   |               |
|---|---------------|
| DNA   | SEQ ID NO: 21 |
| Amino Acid Sequence   | SEQ ID NO: 22 |
| aag ctt ccc aga ccc atg gct gga cct gcc acc cag agc ccc atg aag ctg atg gcc ctg<br>HindIII M A G P A T Q S P M K L M A L 60                   |               |
| -30 -20   |               |
| cag ctg ctg ctg tgg cac agt gca ctc tgg aca gtg cag gaa gcc acc ccc ctg ggc cct 120<br>Q L L L W S A L W T V Q E A T P L G P                  |               |
| -10 -1 1  |               |
| gcc agc tcc ctg ccc cag agc ttc ctg ctc aag tgc tta gag caa gtg agg aag atc cag 180<br>A S S L P Q S F L L K C L E Q V R K I Q                |               |
| 10 20   |               |
| ggc gat ggc gca gcg ctc cag gag aag ctg tgt gcc acc tac aag ctg tgc cac ccc gag 240<br>G D G A A L Q E K L C A T Y K L C H P E                |               |
| 30 40   |               |
| gag ctg gtg ctg ctc gga cac tct ctg ggc atc ccc tgg gct ccc ctg agc agc tgc ccc 300<br>E L V L L G H S L G I P W A P L S S C P                |               |
| 50 60   |               |
| agc cag gcc ctg cag ctg gca ggc tgc ttg agc caa ctc cat agc ggc ctt ttc ctc tac 360<br>S Q A L Q L A G C L S Q L H S G L F L Y                |               |
| 70 80   |               |
| cag ggg ctc ctg cag gcc ctg gaa ggg atc tcc ccc gag ttg ggt ccc acc ttg gac aca 420<br>Q G L L Q A L E G I S P E L G P T L D T                |               |
| 90 100  |               |
| ctg cag ctg gac gtc gcc gac ttt gcc acc acc atc tgg cag cag atg gaa gaa ctg gga 480<br>L Q L D V A D F A T T I W Q Q M E E L G                |               |
| 110 120   |               |
| atg gcc cct gcc ctg cag ccc acc cag ggt gcc atg ccg gcc ttc gcc tct gct ttc cag 540<br>M A P A L Q P T Q G A M P A F A S A F Q                |               |
| 130 140   |               |
| cgc cgg gca gga ggg gtc cta gtt gcc tcc cat ctg cag agc ttc ctg gag gtg tgc tac 600<br>R R A G G V L V A S H L Q S F L E V S Y                |               |
| 150 160   |               |
| cgc gtt cta cgc cac ctt gcc cag ccc gga tcc ggt ggc ggt tcc ggt gga ggc gga agc 660<br>R V L R H L A Q P G S G G G S G G G G S                |               |
| 170 180   |               |
| ggc ggt gga gga gac aaa act cac aca tgc cca ccg tgc cca gca cct gaa gtc ggc 720<br>G G G G S D K T H T C P P C P A P E <u>V</u> <u>A</u>      |               |
| 190 200   |               |
| ggg gga ccg tca gtc ttc ctc ttc ccc cca aaa ccc aag gac acc ctc atg atc tcc cgg 780<br>G G P S V F L F P P K P K D T L M I S R                |               |
| 210 220   |               |
| aca cct gag gtc aca tgc gtg gtg gtg gac gtg agc cac gaa gac cct gag gtc aag ttc 840<br>T P E V T C V V V D V S H E D P E V K F                |               |
| 230 240   |               |
| aac tgg tac gtg gac ggc gtg gag gtg cat aat gcc aag aca aag ccg cgg gag gag cag 900<br>N W Y V D G V E V H N A K T K P R E E Q                |               |
| 250 260   |               |
| tac aac agc acg tac cgg gtg gtc agc gtc ctc acc gtc ctg cac cag gac tgg ctg aat 960<br>Y N S T Y R V V S V L T V L H Q D W L N                |               |
| 270 280   |               |
| ggc aag gag tac aag tgc aag gtc tcc aac aaa gcc ctc cca gcc <u>tcc</u> atc gag aaa acc 1020<br>G K E Y K C K V S N K A L P A <u>S</u> I E K T |               |
| 290 300   |               |
| atc tcc aaa gcc aaa ggg cag ccc cga gaa cca cag gtg tac acc ctg ccc cca tcc cgg 1080<br>I S K A K G Q P R E P Q V Y T L P P S R               |               |
| 310 320   |               |
| gat gag ctg acc aag aac cag gtc agc ctg acc tgc ctg gtc aaa ggc ttc tat ccc agc 1140<br>D E L T K N Q V S L T C L V K G F Y P S               |               |
| 330 340   |               |
| gac atc gcc gtg gag tgg gag agc aat ggg cag ccg gag aac aac tac aag acc acg cct 1200<br>D I A V E W E S N G Q P E N N Y K T T P               |               |
| 350 360   |               |
| ccc gtg ctg gac tcc gac ggc tcc ttc ttc ctc tac agc aag ctc acc gtg gac aag agc 1260<br>P V L D S D G S F F L Y S K L T V D K S               |               |
| 370 380   |               |
| agg tgg cag cag ggg aac gtc ttc tca tgc tcc gtg atg cat gag gct ctg cac aac cac 1320<br>R W Q Q G N V F S C S V M H E A L H N H               |               |
| 390 400   |               |
| tac acg cag aag agc ctc tcc ctg tct ccg ggt aaa tga gaa ttc 1365<br>Y T Q K S L S L S P G K EcoRI   |               |
| 410 417   |               |